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鲢科鱼类细胞色素 b 基因片段的序列测定及其系统发育的初步研究

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何舜平^{①②} 陈永久^② 张亚平^②

Q959.468

(①中国科学院水生生物研究所淡水鱼类进化生物学和生物地理学研究室 武汉 430072)

(②中国科学院昆明动物研究所细胞与分子进化开放研究实验室 昆明 650223)

摘 要 从 9 种鲢科鱼类的福尔马林标本中获得了 333 bp 的细胞色素 b 基因片段的序列。这 9 个种分别代表鲢科鱼类的 8 个属。333 bp 的 DNA 序列经 MUST 软件排序后, 有 101 个变异位点, 其中有 39 个信息位点。序列在成对物种间的距离为 8~48。平均遗传距离为 2.4%~14.4%。简约分析产生了最大简约系统树, 其步长是 162 (CI=0.735, RI=0.494)。在该系统树上, *Bagarius* 是

最原始的属, 并与所有其他的物种形成姊妹群。其余 8 个属形成一个单系类群并分为二个姊妹群。尽管在形态上具有 13 个离征, 但在分子系统树上, 鲤科鱼类并未形成一个单系类群。可能的原因是 333 bp 序列中的星系信息位点太少; 另外单从福尔马林浸制标本获得的 DNA 序列的可靠性尚有待进一步验证。

关键词 mtDNA, 细胞色素 b, 系统发育, 福尔马林浸制标本, 鲢科鱼类
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